

PATENT

Client-Matter No.: 66661-036 (P-IS 4988)

## IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of Ideker and Hood

Serial No: 09/993,312

Filed: November 13, 2001

For: MULTIPARAMETER INTEGRATION)

METHODS FOR THE ANALYSIS OF BIOLOGICAL NETWORKS

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Confirmation No.: 5632 Group Art Unit: 1631 Examiner: C. Smith

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#### PRELIMINARY AMENDMENT

This Preliminary Amendment is filed in connection with the above-identified patent application, which claims the benefit of priority to U.S. Provisional Application No. 60/248,257, filed November 14, 2000, and U.S. Provisional Application No. 60/266,038, filed February 2, 2001. Entry of the amendment and ed. consideration of the following remarks is respectfully requested.

#### **AMENDMENTS**

#### In the specification:

Please replace Table 1 on pages 110 to 113, with the following replacement Table 1, which shows changes made relative to the immediate prior version by underlining of added text.

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Table 1: Gal4p binding-site predictions

Cluster	Gene	Upstream	Core	Matrix	Sequence
		position	similarity	similarity	
	i	(+/-	score	score	
		strand)			
1	GAL1	456(+)	1.000	0.840	GTACGGATTAG
	Ì				AAGCCGCCGAG
					С
	,				(SEQ ID
<u></u>					NO:1)
1	GAL1	437 (+)	1.000	0.880	GAGCGGGCGAC
				·	AGCCCTCCGAC
<u> </u>		į			G
	:				(SEQ ID
					<u>NO:2)</u>
1	GAL1	419(+)	0.875	0.920	CGACGGAAGAC
	:				TCTCCTCCGTG
					С
					(SEQ ID
					<u>NO:3)</u>
1	GAL1	355 (+)	1.000	0.860	CCTCGCGCCGC
					ACTGCTCCGAA
:					С
					(SEQ ID
					NO:4)
1	GAL10	336(-)	1.000	0.860	CCTCGCGCCGA
					ACTGCTCCGAA
			·		С
					(SEQ ID
					NO:4)

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	7 —	1	<del></del>	T	T
1	GAL10	272 (-)	0.875	0.920	CGACGGAAGAC
					TCTCCTCCGTG
					С
					(SEQ ID
					NO:5)
1	GAL10	254 (-)	1.000	0.880	GAGCGGGCGAC
					AGCCCTCCGAC
					G
			1		(SEQ ID
					NO:6)
1	GAL10	235 (-)	1.000	0.840	GTACGGATTAG
			ļ		AAGCCGCCGAG
	i				C (SEQ ID
					NO:7)
1	GAL7	282(-)	1000	0.870	CTTCGCTCAAC
					AGTGCTCCGAA
					G (SEQ ID
					NO:8)
1	GAL7	195(-)	1.000	0.924	TCACGGTCAAC
					AGTTGTCCGAG
					C (SEQ ID
					NO:9)
1	GAL7	188 (+)	1.000	0.848	CAACTGTTGAC
					CGTGATCCGAA
					G (SEO ID
					<u>NO:10)</u>
1	GCY1	372 (-)	1.000	0.828	CCCCGGAATAG
					TCTGCCCCGAT
					T (SEQ ID
					NO:11)

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1	PCL10	235(-)	1.000	0.905	GATCGGTGCAA TATACTCCGAG C (SEQ ID
					NO:12)
1	GAL2	533(-)	1.000	0.818	TTCCGGAAGGA AGCTTTCCGAA T (SEQ ID NO:13)
1	GAL2	419(+)	0.875	0.903	CACCGGCGGTC TTTCGTCCGTG C (SEO ID NO:14)
1	GAL2	400(-)	0.800	0.825	GAACGGCGCAG ATATCTCCGCA C (SEO ID NO:15)
1	GAL2	336(+)	1.000	0.867	TATCGGGGCGG ATCACTCCGASC C (SEO ID NO:16)
1	GAL2	331 (+)	1.000	0.855	GGGCGGATCAC TCCGAACCGAG A (SEO ID NO:17)
1	YEL057C	417 (-)	0.875	0.811	CCCCGGACGGC AGCCGCCGTC C (SEQ ID NO:18)

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		1		<del></del>	<del></del>
1	YGR090W	381(-)	0.875	0.819	CAACGGCATGC
					AGCGAGCCGTA
					G (SEQ ID
					NO:19)
1	YPL066W	101(+)	1.000	0.856	TCACGGTCATC
					ACTGCTCCGAC
					A <u>(SEQ ID</u>
					NO:20)
2	GAL3	291(-)	1.000	0.938	GTTCGGCACAC
ļ					AGTGGACCGAA
		į			C (SEQ ID
					NO:21)
2	GAL80	175(+)	1.000	0.952	TACCGGCGCAC
					TCTCGCCCGAA
					C (SEO ID
					NO:22)
2	YPR194C	624 (-)	1.000	0.814	CGTCGGACAGC
					AACCCCCGAT
					T <u>(SEQ ID</u>
77.44		-			NO:23)
2	YMR318C	239 (+)	1.000	0.830	GTCCGGTCCGT
					CCTTGACCGAA
					G <u>(SEQ ID</u>
					NO:24)
11	RPA49	249(+)	1.000	0.804	GACCGGACACC
					TAATCACCGAC
					G (SEQ ID
					NO:25)

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11	YLR201C	143(-)	1.000	0.812	CTTCCGCCTAA
		143( )	1.000	0.812	TATAGTCCGAA
					1
					A (SEQ ID
1.5		50.4 \			NO:26)
15	YJL045W	524(-)	1.000	0.804	CGACGGGGAAT
					TGAACCCCGAT
					C (SEQ ID
					NO:27)
15	MBR1	313(+)	0.800	0.811	GAGCGGCTCCC
i I					CTTTCCC
					CGGAA <u>(SEQ</u>
					<u>ID NO:28)</u>
15	MRK1	268(-)	0.725	0.828	CATCGGACGA
					CTTTGCTCCC
					AGG (SEQ
					ID NO:29)
15	YMR031	305(-)	1.000	0.801	TTTTGGGTAA
	С				CAGCGGACCG
					AAG <u>(SEQ</u>
					<u>ID NO:30)</u>
16	ICL1	407(+)	1.000	0.810	CCCAGGTTTC
					CATTCATCCG
					AGC <u>(SEQ</u>
					<u>ID NO:31)</u>
16	YLR164	243 (+)	1.000	0.836	GATTGGAGTA
	W				CCCTTATCCG
					AAG <u>(SEQ</u>
					ID NO:32)
	<u>-</u>	l			

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16	YIL057	192(+)	0.875	0.867	CGGCGGTTGG
	С				CAATCGTCCG
					TAT <u>(SEQ</u>
					<u>ID NO:33)</u>

# Regarding the amendment

The amendment to the specification was made to insert sequence identification numbers and accordingly adds no new matter. Therefore, entry of the amendment is respectfully requested.

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### CONCLUSION

No fee is deemed necessary in connection with the filing of this Communication. However, if any fee is required, authorization is given to charge the amount of this fee to Deposit Account No. 50-2624. The Examiner is invited to call the undersigned agent or Cathryn Campbell if there are any questions relating to this application.

Respectfully submitted,

September 25, 2003

Date

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